

09/486094  
428 Rec'd PCT/PTO 18 FEB 2000

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SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT:

- 5 (A) NAME: RHONE-POULENC AGROCHIMIE  
(B) STREET: 14-20 Rue Pierre BAIZET  
(C) TOWN: LYONS  
(E) COUNTRY: France  
(F) POST CODE: 69009

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- (ii) TITLE OF THE INVENTION: Gene coding for  
androctonine, vector containing it and  
disease-resistant transformed plants obtained

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- (iii) NUMBER OF SEQUENCES: 11

(vi) COMPUTER-READABLE FORM:

- (A) MEDIUM TYPE: Floppy disk  
(B) COMPUTER: IBM PC compatible  
20 (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: PatentIn Release #1.0,  
Version #1.30 (EPO)

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(2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 110 base pairs  
(B) TYPE: nucleotide

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- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA

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(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..75

10 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

```
AGG TCC GTG TGC AGG CAG ATC AAG ATC TGC AGG AGG AGG GGT GGT TGC 48
Arg Ser Val Cys Arg Gln Ile Lys Ile Cys Arg Arg Arg Gly Gly Cys
1           5           10           15
15 TAC TAC AAG TGC ACT AAC AGG CCA TAC TGAGCTCGGC GAGGCGAAGC 95
Tyr Tyr Lys Cys Thr Asn Arg Pro Tyr
20           25
TGTCGACGGG TCCGG 110
```

20 (2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 106 base pairs
- (B) TYPE: nucleotide
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: DNA

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(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 12..101

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

GGGTGACGC C ATG GGT TTC GTG CTT TTC TCT CAG CTT CCA TCT TTC CTT 30

Met Gly Phe Val Leu Phe Ser Gln Leu Pro Ser Phe Leu

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CTT GTG TCT ACT CTT CTT CTT TTC CTT GTG ATC TCT CAC TCT TGC CGT 30

Leu Val Ser Thr Leu Leu Leu Phe Leu Val Ile Ser His Ser Cys Arg

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106

15 GCC GCGCA

Ala

30

(2) INFORMATION FOR SEQ ID NO: 3:

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(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 211 base pairs

(B) TYPE: nucleotide

(C) STRANDEDNESS: double

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(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA

007120-46098460

4

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 12..176

5 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

GGTCGACGC C ATG GGT TTC GTG CTT TTC TCT CAG CTT CCA TCT TTC CTT 50

Met Gly Phe Val Leu Phe Ser Gln Leu Pro Ser Phe Leu

1 5 10

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CTT GTG TCT ACT CTT CTT CTT TTC CTT GTG ATC TCT CAC TCT TGC CGT 98

Leu Val Ser Thr Leu Leu Leu Phe Leu Val Ile Ser His Ser Cys Arg

15 20 25

15 GCC AGG TCC GTG TGC AGG CAG ATC AAG ATC TGC AGG AGG AGG GGT GGT 146

Ala Arg Ser Val Cys Arg Gln Ile Lys Ile Cys Arg Arg Arg Gly Gly

30 35 40 45

TGC TAC TAC AAG TGC ACT AAC AGG CCA TAC TGAGCTCGGC GAGGCGAAGC 196

20 Cys Tyr Tyr Lys Cys Thr Asn Arg Pro Tyr

50 55

TATGACCGA TCCGG 211

25 (2) INFORMATION FOR SEQ ID NO: 4:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 75 base pairs

(B) TYPE: nucleotide

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(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

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(A) DESCRIPTION: /desc = "synthetic  
oligonucleotide 1"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

10 GCGTCGAGGC GATGGGTTTC GTCCTTTTCT CTCAGCTTCC ATCTTTCCTT CTGCTGCTCA 60  
CTCTTCTTCT TTTC 75

(2) INFORMATION FOR SEQ ID NO: 5:

(1) SEQUENCE CHARACTERISTICS:

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(A) LENGTH: 72 base pairs  
(B) TYPE: nucleotide  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic  
oligonucleotide 2"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

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TCGCCGCGCAC GCGAAGAGTA AGAGATCACA AGGAAAGCAA GAAGAGTAGA CACAAGAAGG 60  
AAAGATGGAA GC 72

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(2) INFORMATION FOR SEQ ID NO: 6:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 44 base pairs

(B) TYPE: nucleotide

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic  
oligonucleotide 3"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

AGGTCCCTGT GCAGGCAGAT CAGATCTGC ADGAGGAGGG GTGG

(2) INFORMATION FOR SEQ ID NO: 7:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 97 base pairs

(B) TYPE: nucleotide

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic  
oligonucleotide 4"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

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CCGGATCCGT CGACACGTTT GCCTCGCCGA GTCAGTATG GCTGTATTG GCACTTGTAG 60  
TAGCAACCAC CCTCTCTCTT GCAGATCTTG ATCTGCC 97

(2) INFORMATION FOR SEQ ID NO: 8:

- 5 (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 85 base pairs
  - (B) TYPE: nucleotide
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

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- (ii) MOLECULE TYPE: other nucleic acid
- (A) DESCRIPTION: /desc = "synthetic  
oligonucleotide 5"

15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

AGGGCCCCCT AGGCTTTAAA CCGCCAGTCA GGCAGATTC GAGCTCGGTA CCGGGGGATC 60  
CTCTAGAGTC GACCTGCAGG CATGC 85

20 (2) INFORMATION FOR SEQ ID NO: 9:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 66 base pairs
  - (B) TYPE: nucleotide
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

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- (ii) MOLECULE TYPE: other nucleic acid
- (A) DESCRIPTION: /desc = "synthetic"

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oligonucleotide 6"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

5 CCTCGAACCA GGCTCGAGGG CCGGCCCTTAA TTAAGGCTT GCATGCCCTGC AGGTGCACTC 60  
TAGAGG 66

(2) INFORMATION FOR SEQ ID NO: 10:

(i) SEQUENCE CHARACTERISTICS:

10 (A) LENGTH: 93 base pairs  
(B) TYPE: nucleotide  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

15 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic  
oligonucleotide 7"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

20 CCGGCCAGTC AGGCCCACT TAATTAAAGTT TAAACGCCGC CCGGCCCGGC CTAGGTGTGT 60  
GCTCGAGGGC CCAACCTCAG TACCTGGTTC AGG 93

(2) INFORMATION FOR SEQ ID NO: 11:

25 (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 93 base pairs  
(B) TYPE: nucleotide  
(C) STRANDEDNESS: single

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(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic  
oligonucleotide 8"

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

CCGGCCTGAA CCAGGTACTG AGTTGGGCC CCGAGCACA CACCTAGCG CGCCGGGGCC 60  
10 GCCTTTAAAC TTAATTAAGT GTGGCCTGAC TGG 93

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